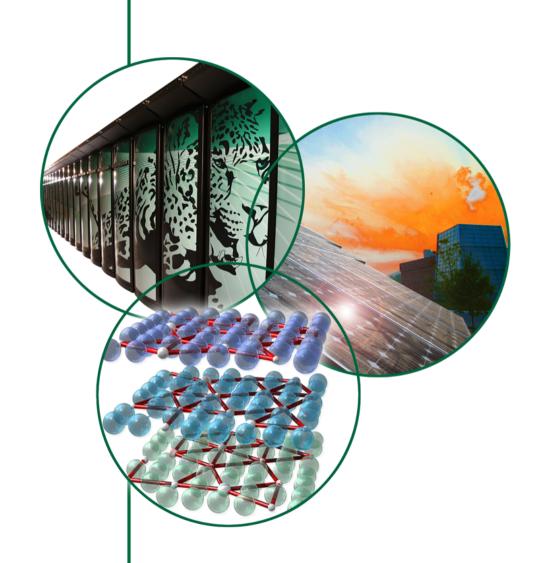
## Agent Design and Performance

Valmor de Almeida Benjamin Hay

**NEAMS PI Meeting** 

L'Enfant Plaza Hotel, Washington DC 18-20 October 2010





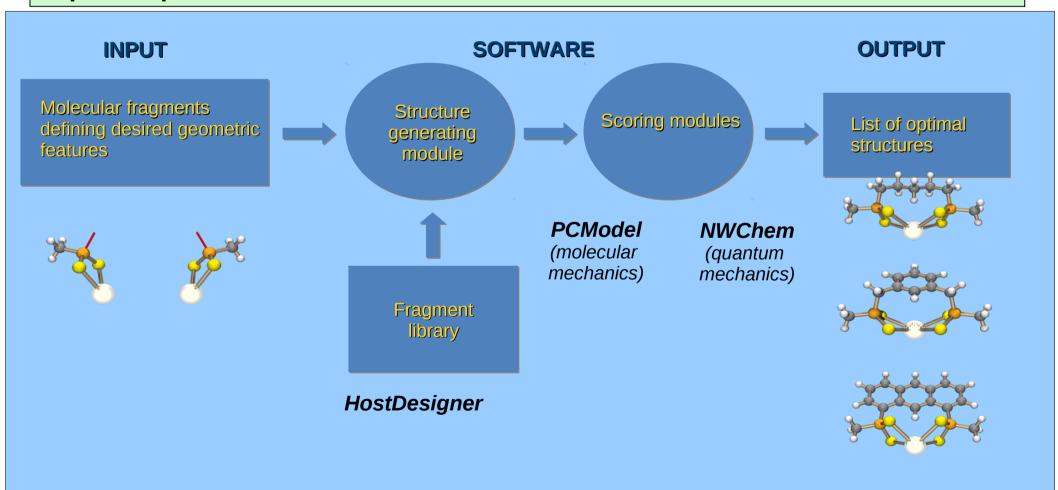


## SafeSeps IPSC Challenge Problem: Agent Design to Process Design

- ➤ **Description:** Develop molecular design software to predict performance of sequestering agents for applications in separation of specific radionuclides from nuclear fuel reprocessing plant streams:
  - Improved selectivity, binding affinity, and aqueous insolubility
  - Improved ion-exchange
  - Improved interfacial transport behavior
- ➤ **Impact:** *NE R&D Objective Three*. Highly selective separating agents will be needed to meet the needs associated with future fuel cycles and responsibly disposing of nuclear waste:
  - Save significant cost in agent identification, evaluation and deployment.
  - Key need of Separations and Waste Forms Technical Area of the Fuel Cycle R&D Program (Minor Actinide Sigma Team).

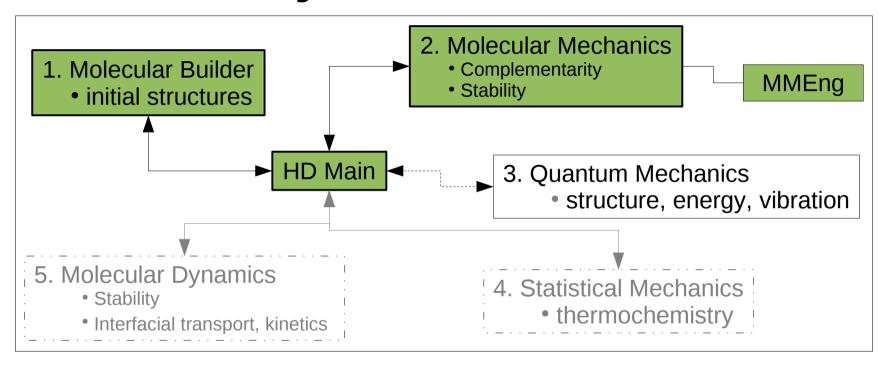
# New ligands for selective complexation of metal ions (HostDesigner)

Computer-aided, structure-based design of radionuclide sequestering agents is being used to guide experimental programs. Increase in fragment library size and need for more accurate scoring motivates adapting this system for supercomputers.



#### **Activity:**

## Develop molecular-level codes for ligand design and performance analysis

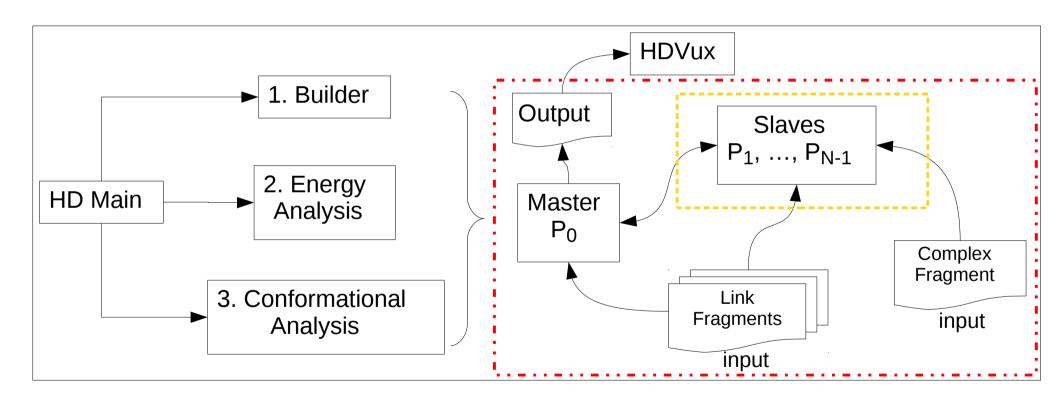


#### FY10 Tasks

- Modifications of HostDesigner code
  - Reorganization to allow for coupling of new modules
  - ✓ Parallelization of molecular mechanics modules
- ✓ Identification of coupling with quantum mechanics libraries
- ✓ Testing

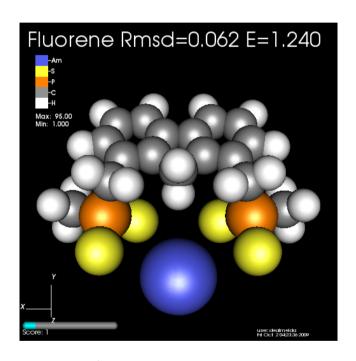
### **Progress on Parallelization**

- ➤ Parallelization of Host-Designer active items
  - ➤ Master-slave approach for balancing the computational work
  - ➤ Planning parallel IO



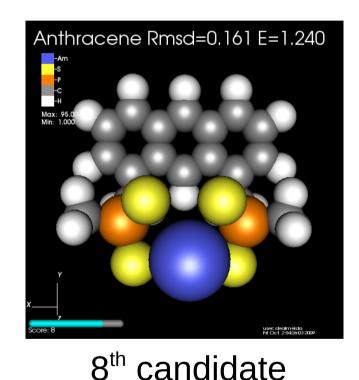
## Verification against sequential code results

>Americium dimethylphosphinate chelate complex fragment



asymmetric

candidates



1<sup>st</sup> candidate

- ➤ Parallel results match sequential code
  - Ligand suggested for synthesis by Minor Actinides Sigma Team



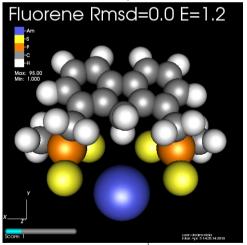
#### **Testing:**

#### **Americium ligand (Minor Actinides Sigma Team)**

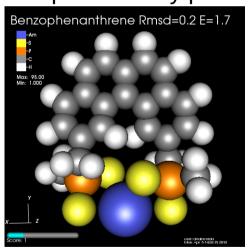
Library – 8 266 Links				
# Proc's	Build (2 188 candidates)	Complementarity (2 188 candidates)	Stability (1 024 candidates)	
1024	2.5 min	2.7 min	98.8 min	

- > Previous simulation capability: limited MM analysis on 10 to 20 ligands (~days wall-clock time)
- > Current capability: MM conformer search on > 1000 ligands in ~1.5 hours wall-clock time

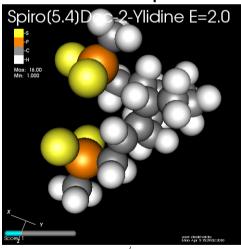
#### Build phase



#### Complementarity phase



#### Conformer phase



top candidates in each phase

- > Modularized and integrated molecular mechanics module tested on Cray-XT5 parallel machine.
- New MPI version of the code to be used in FY10 Minor Actinides Sigma Team work.

## **Cluster Computing Testing**

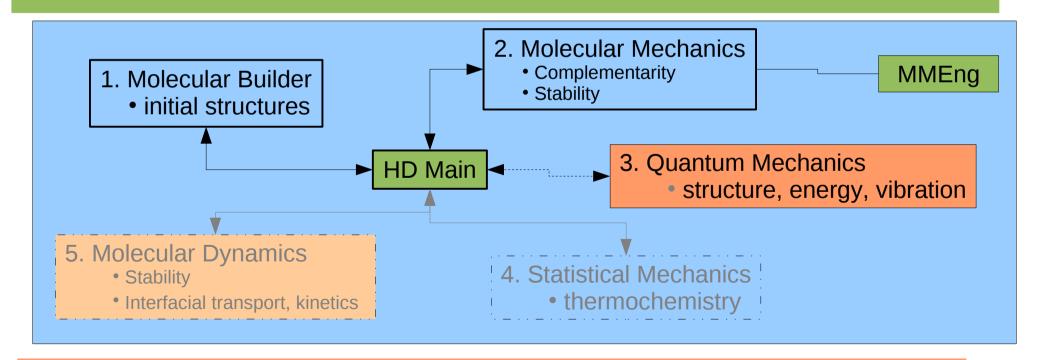
Library – 8 266 Links				
# Proc's	Build (1 094 candidates)	Complementarity (1 094 candidates)	Stability (256 candidates)	
256	2.0 min	1.0 min	74.0 min	

- > INL CAMS cluster Helios
- Demonstrated portability on common linux cluster environment
- > Potential source of CPU cycles for production runs by FCR&D users
- Would like access to more powerful machines for QM calculations

#### **Outlook**

#### Status, issues, and FY 11 work

- > Sigma Team users now equipped with more powerful design capability needed in FY10
  - Experimental testing of synthesized molecules indicated promising extraction performance
  - But poor resistance to chemical environment
- > Molecular mechanics module will require further development:
  - Potential rewrite of MMEng code or replacement by existing MM engine
- Module integration may benefit from software architecture analysis or use of a framework



- Quantum Mechanics module will be integrated next (NWChem parallel library)
  - Greatest impact of massively parallel computing to agent design and performance